



FIG. 2

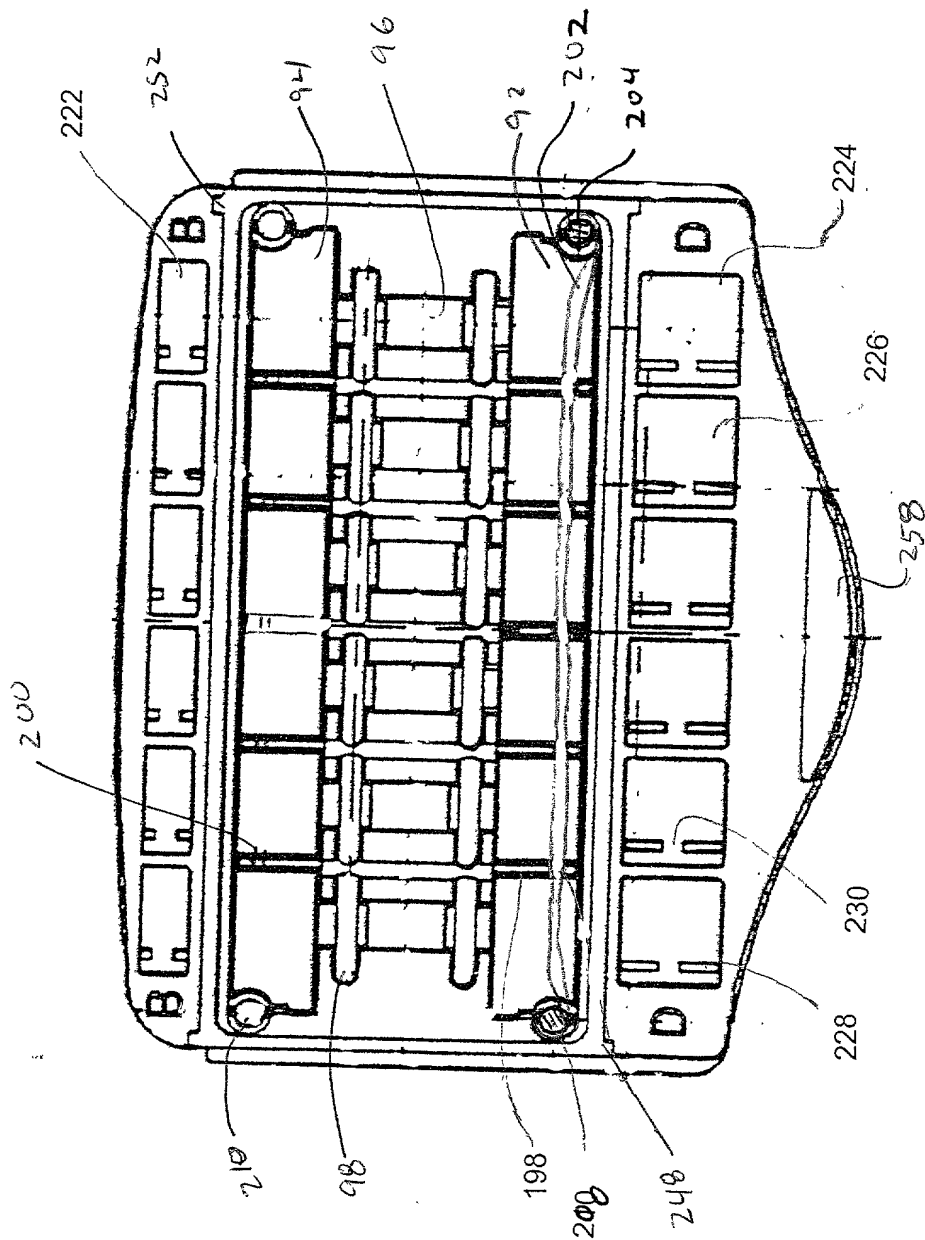
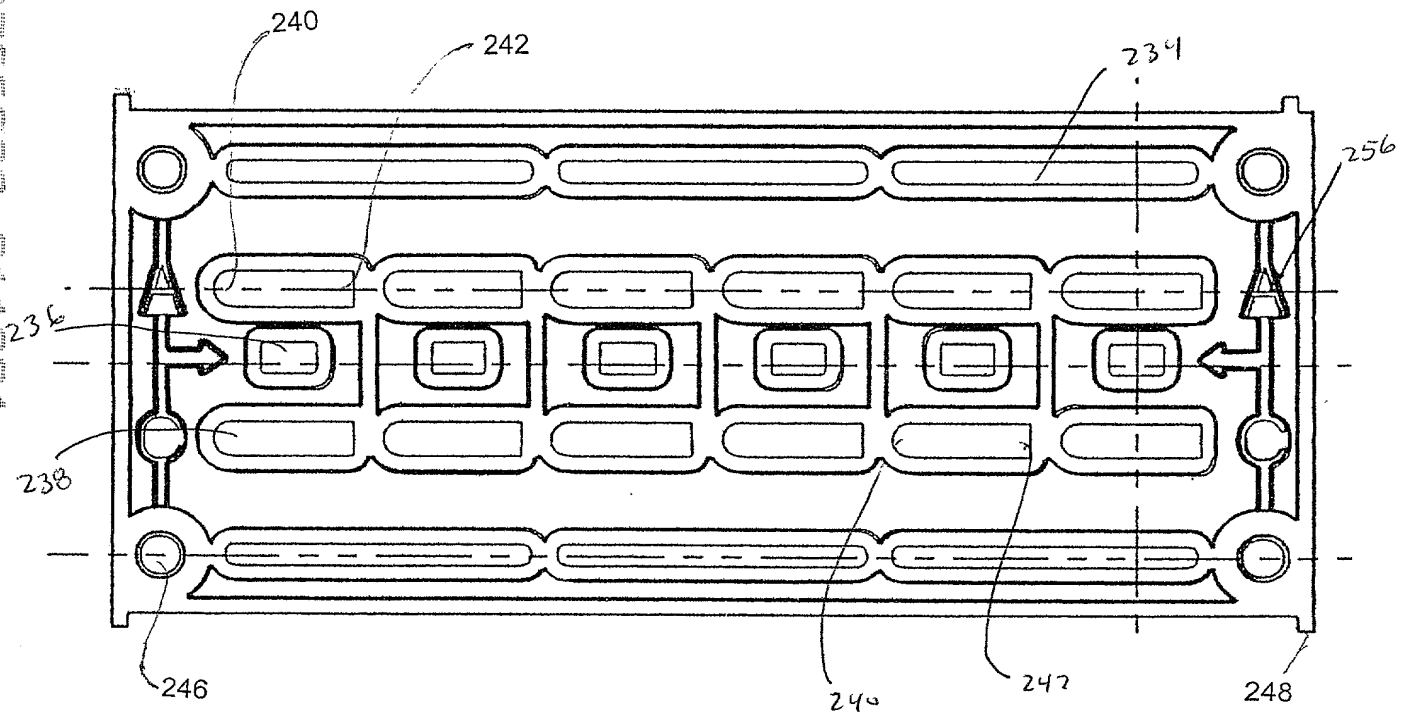
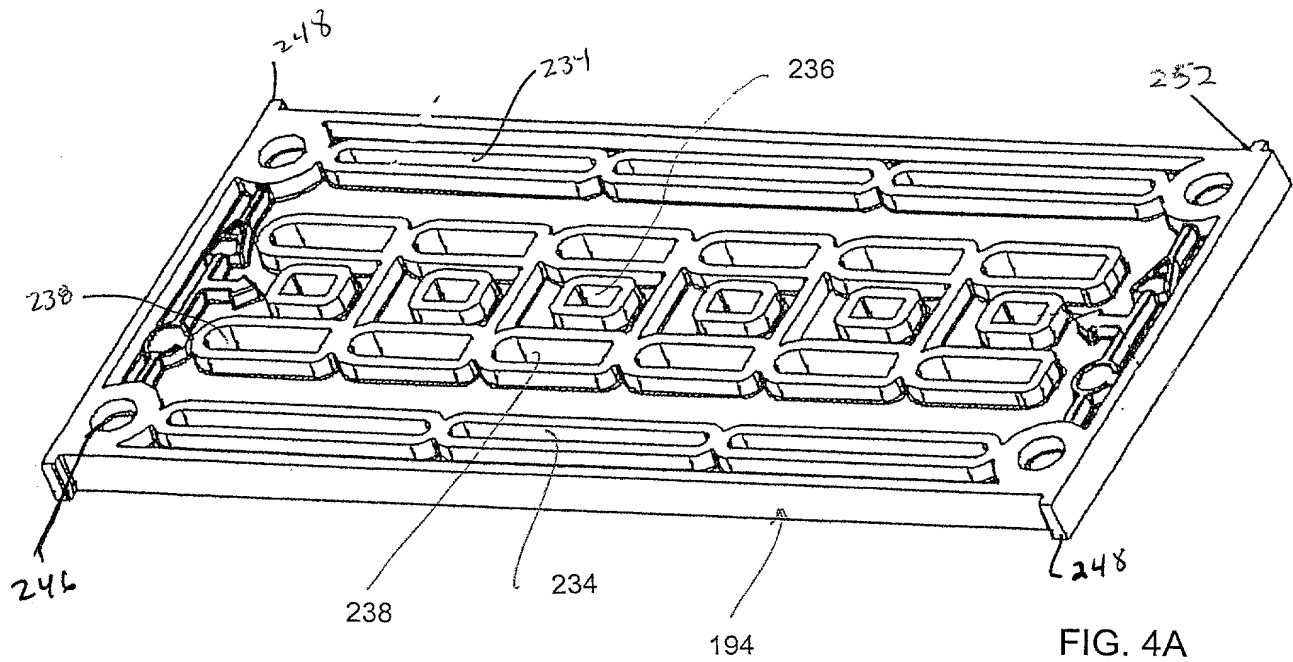


FIG. 3



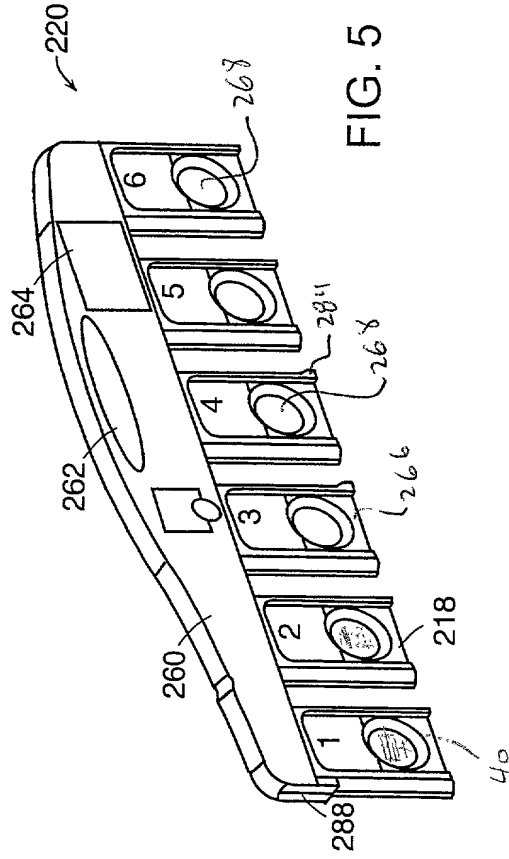


FIG. 5

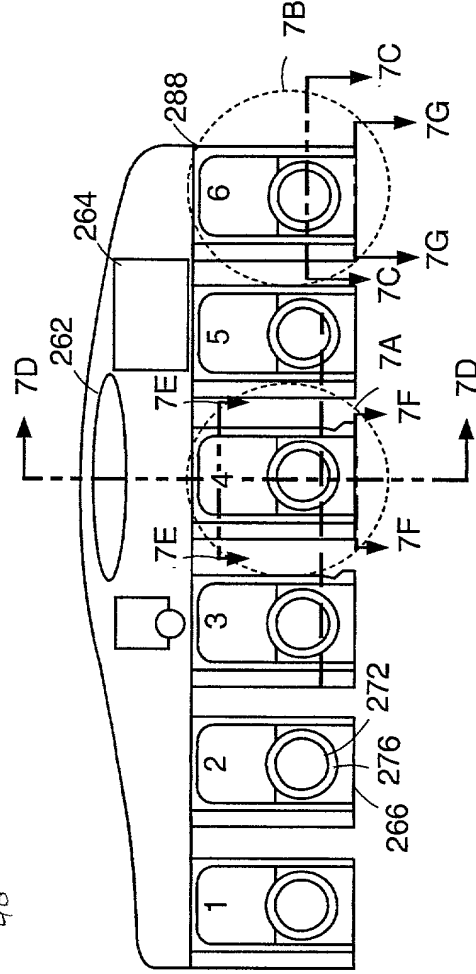


FIG. 6A

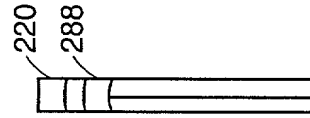


FIG. 6B

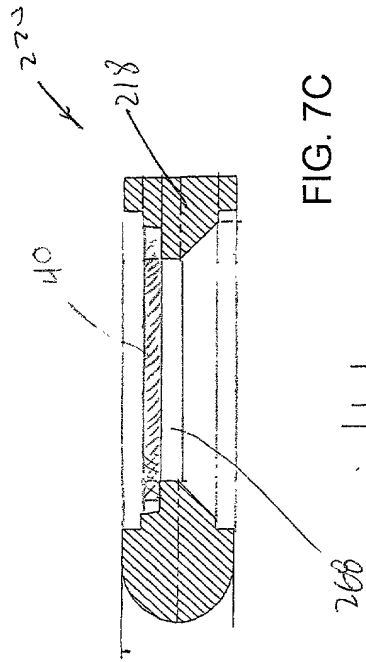


FIG. 7C

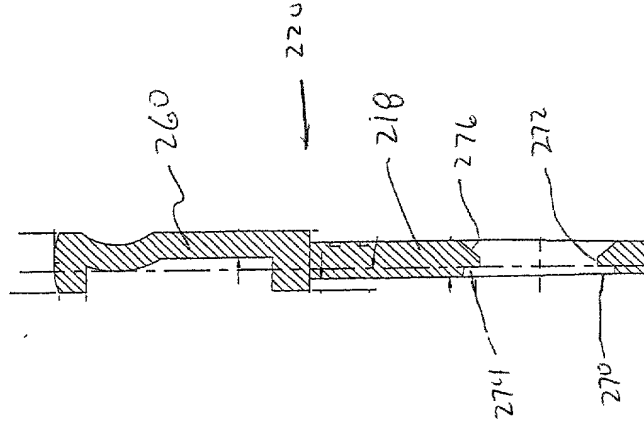


FIG. 7D

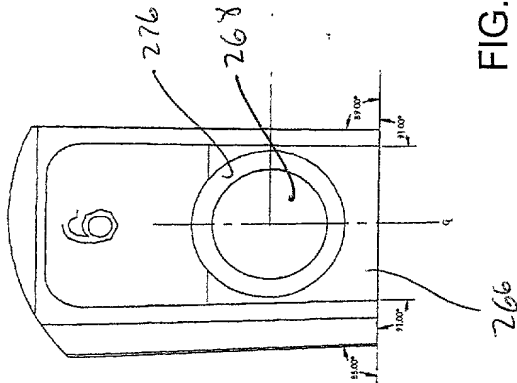


FIG. 7B

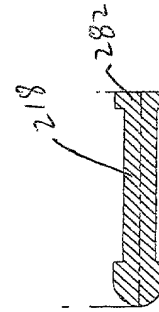


FIG. 7G

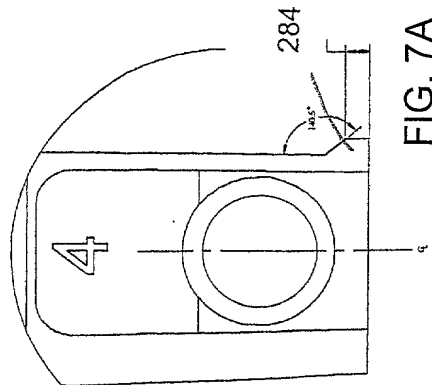


FIG. 7A

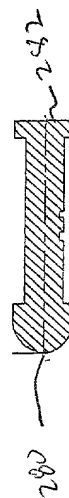


FIG. 7E

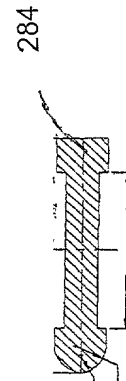
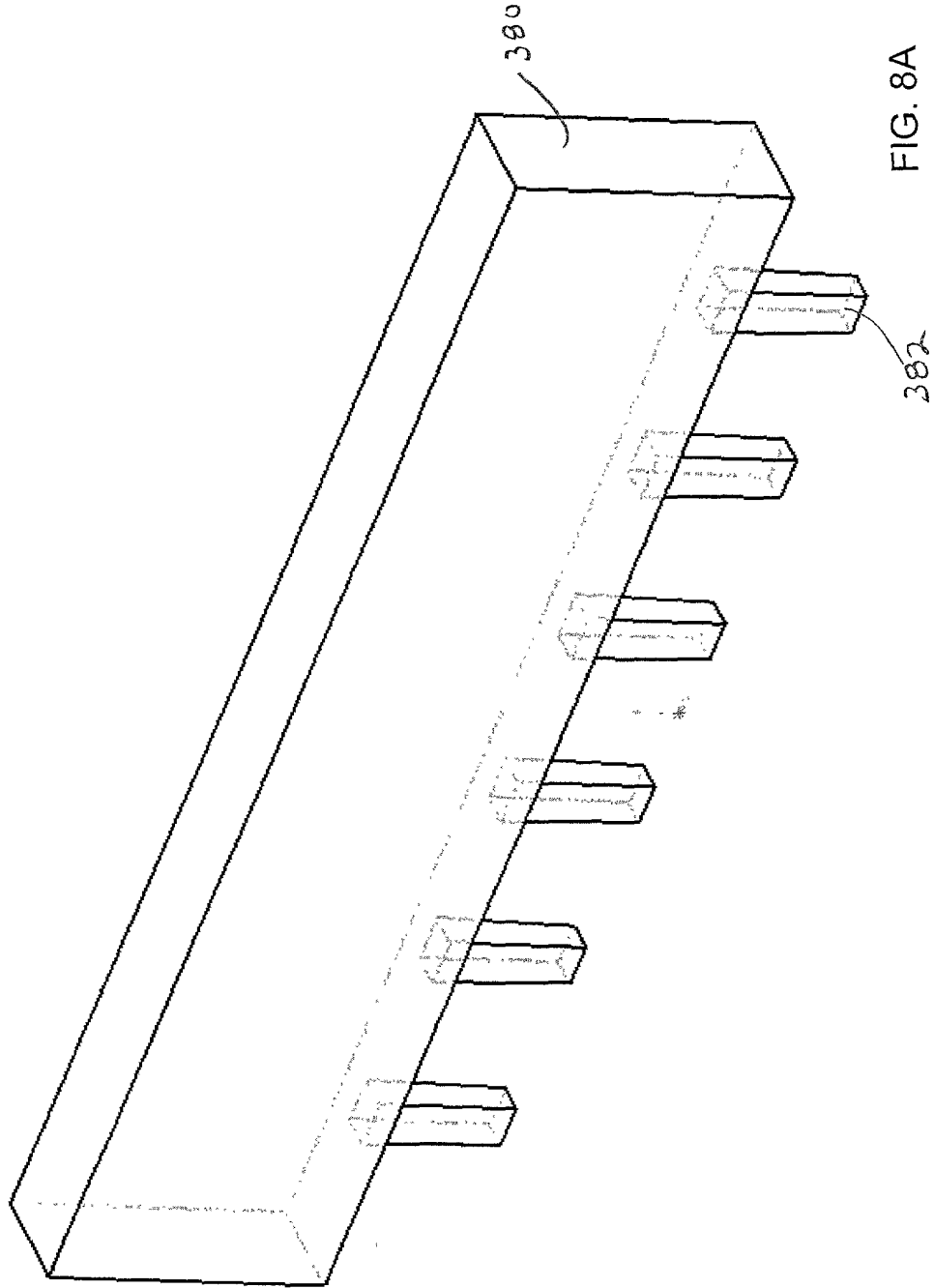


FIG. 7F



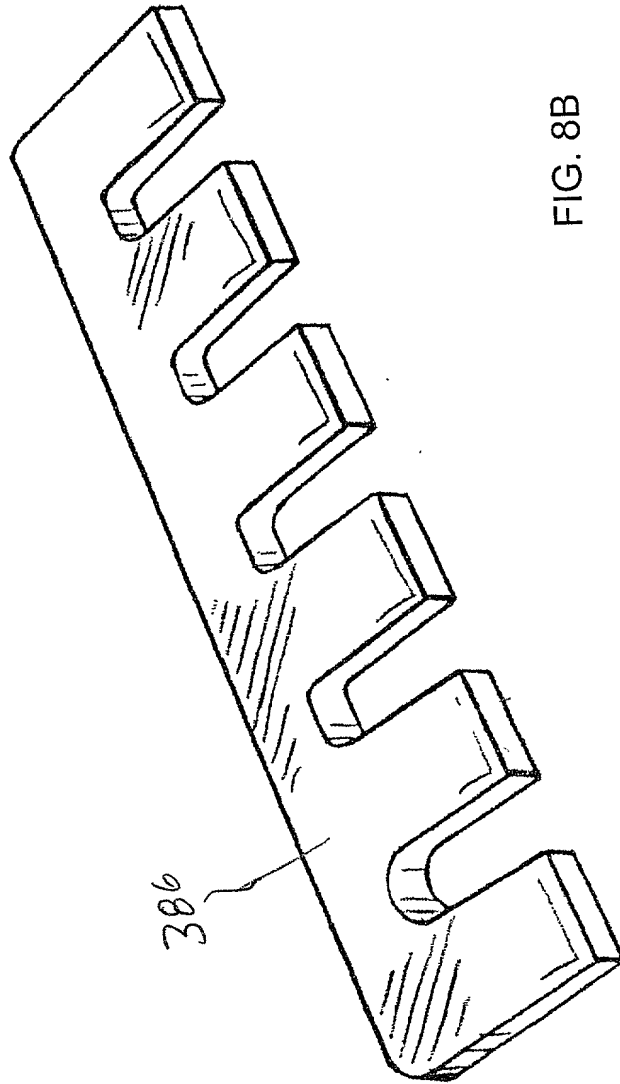
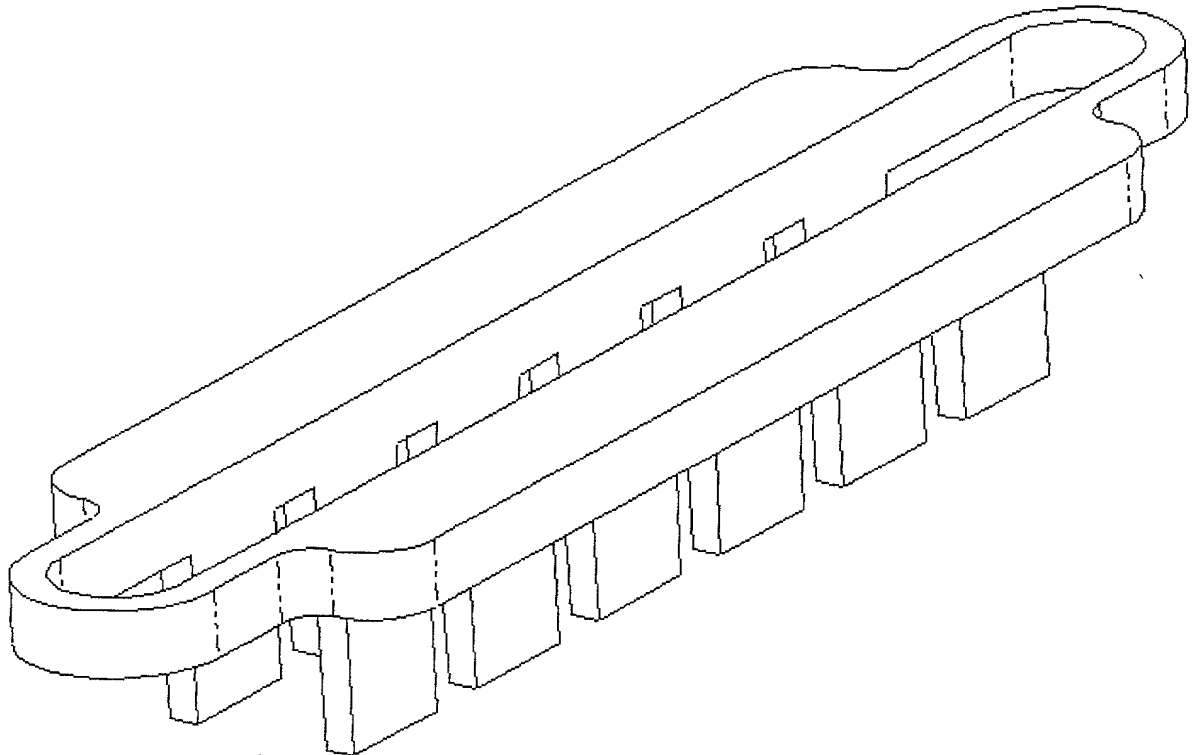


FIG. 8B

FIG. 8B



390

FIG. 8C

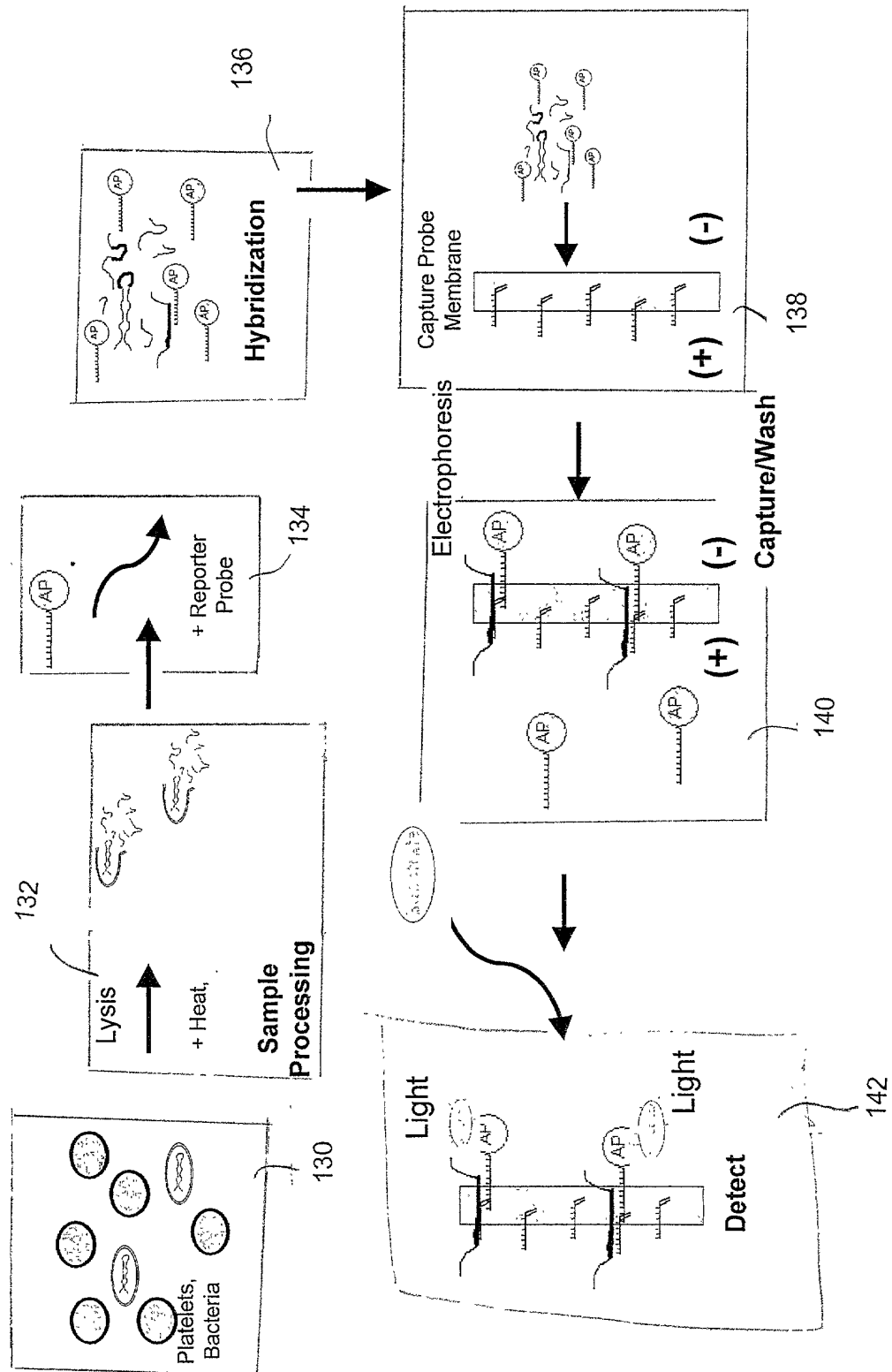


FIG. 9A

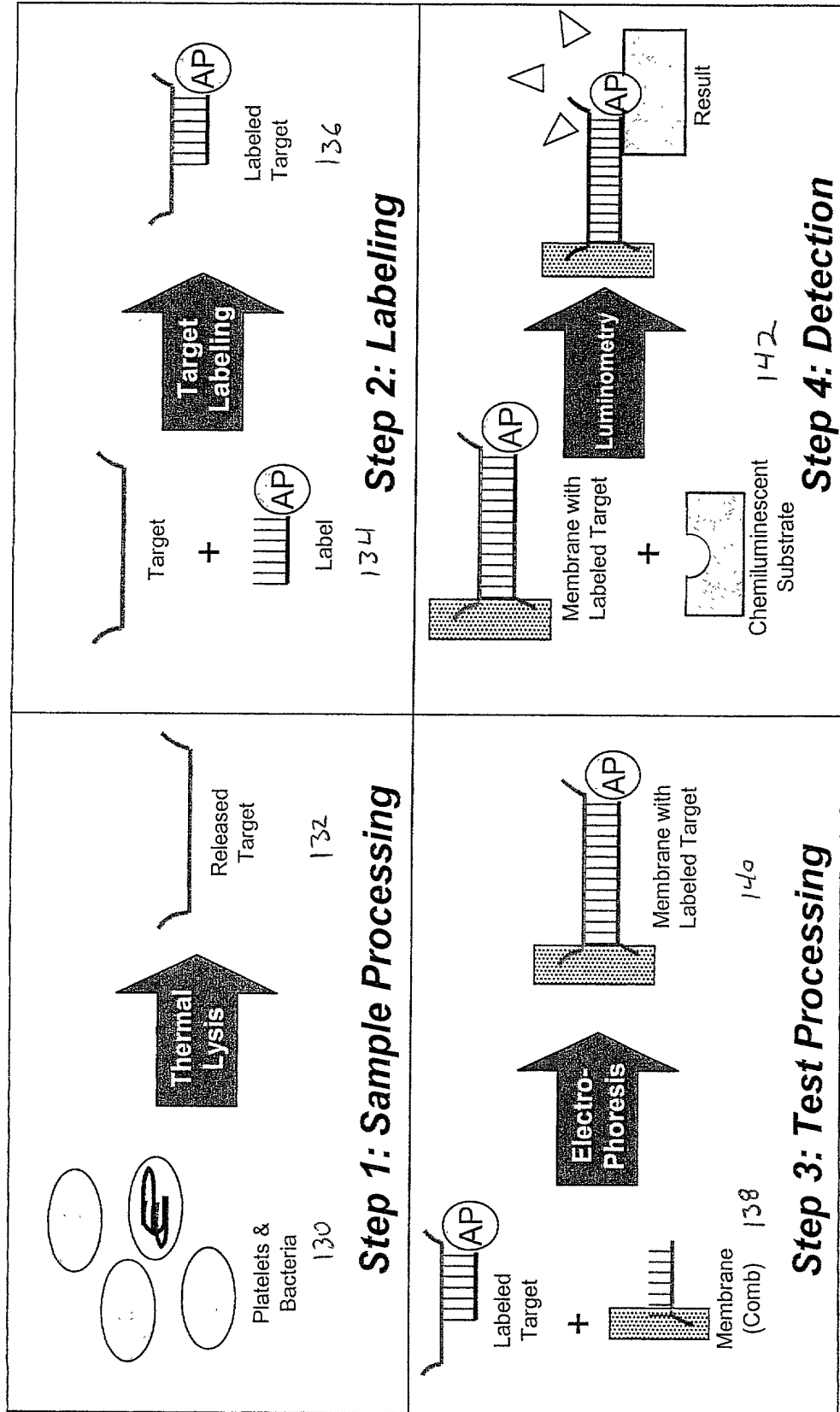


FIG. 9B

Staphylococcus species (* all unfinished)

staph_epi : CATGACCATGTCAGGTCTCTGACCGGAAGCAGCCATTAAGTGGATGCTGATA : 198
 Staph_prob : -----ACATGTCAGGTCTCTGACCGGAAGCAGCC----- : 26
 saureus_co : CATGACCATGTCAGGTCTCTGACCGGAAGCAGCCATTAAGTGGATGCTGATA : 200
 catgaacCATGTCAGGTCTCTGACCGGAAGCAGCattaagtggatc tcata

Streptococcus species (* all unfinished)

Streppyo_ : CTGTGCGGGAAGTGGGTTCAGGGGAGGAATCCAGCAGCCCTAAAGCC : 220
 Streppneu : GTTCTGCGGGAAGCCTGGGTTCAGGGGAGGAATCCAGCAGCCCTAAAGCC : 92
 strep_targ : -----ATGGGTTCAGGGGAGGAATCCAGCAGCC----- : 26
 Strep_Equi : CTGTGCGGGAAGTGGGTTCAGGGGAGGAATCCAGCAGCCCTAAAGCC : 135
 strepmutan : GCTTTCGCGGGAAGCCTGGGTTCAGGGGAGGAATCCAGCAGCCCTAAAGCC : 221
 t tgcgtgaag GGGTCAGGGGAGGAATCCAGCAGCC taagcg

Enterobacteriaceae, Pseudomonas aeruginosa, Bacillus cereus

* klebpneu : CCGAAGCCTAGTCTCTGTTTACCAAGTCAGGTCCCGGAAGGAAGCAGCCCA-GACCAACGACGTCGT : 115
 * Styphimurw : CCGAAGCCTAGTCTCTGTTTACCAAGTCAGGTCCCGGAAGGAAGCAGCCCA-AGGCAACATGACGGCTGT : 84
 ecoli_comp : CCGAAGCCTAGTCTCTGTTTACCAAGTCAGGTCCCGGAAGGAAGCAGCCCA-AGGCAACATGACGGCTGT : 84
 Ecoli_prob : -----ACAGTCAGGTCCCGGAAGGAAGCAGCC----- : 26
 pseud aerug : CCGAAGCCTAGTCTCTGTTTACCAAGTCAGGTCCCGGAAGGAAGCAGCCCA-CACCGCGGAACATCCTGT : 84
 Bacillus_c : CCGAAGCCTAGTCTCTGTTTACCAAGTCAGGTCCCGGAAGGAAGCAGCCCAATAACGGCTCTTCTCCTGT : 194
 cgcaacg c gt acC gGTCAGGTCCCGGAAGGAAGCAGC a gc g gtgt

* unfinished sequence from genome centers

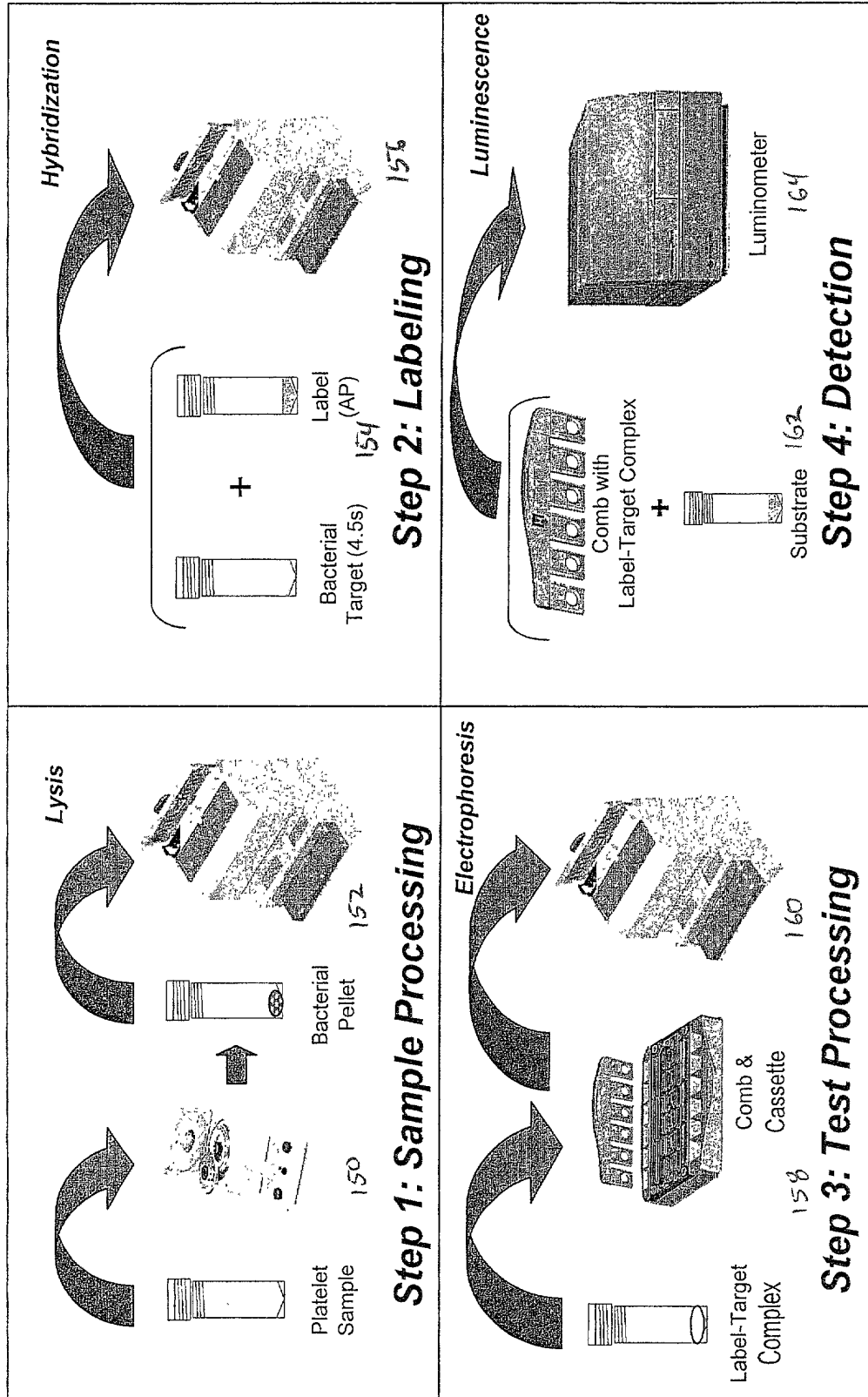
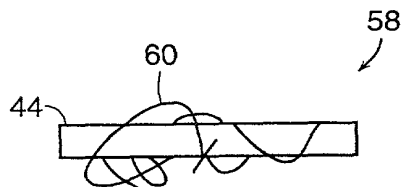
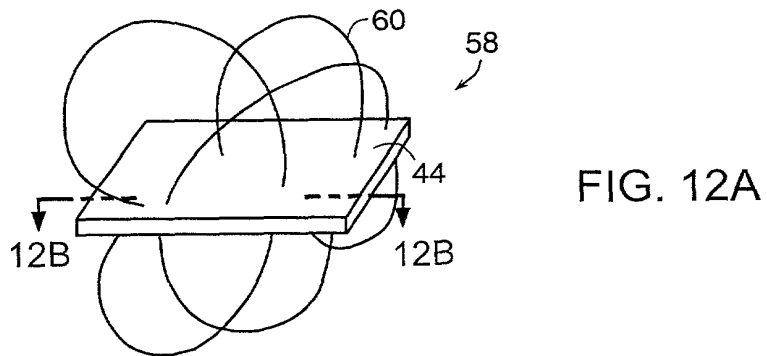
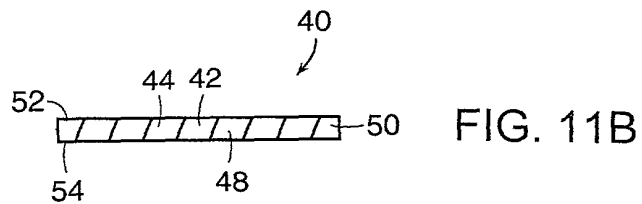
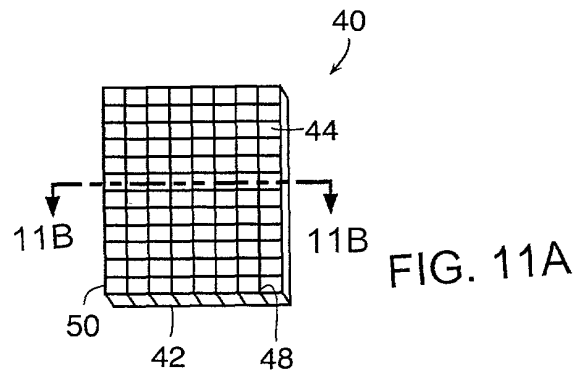


FIG. 10



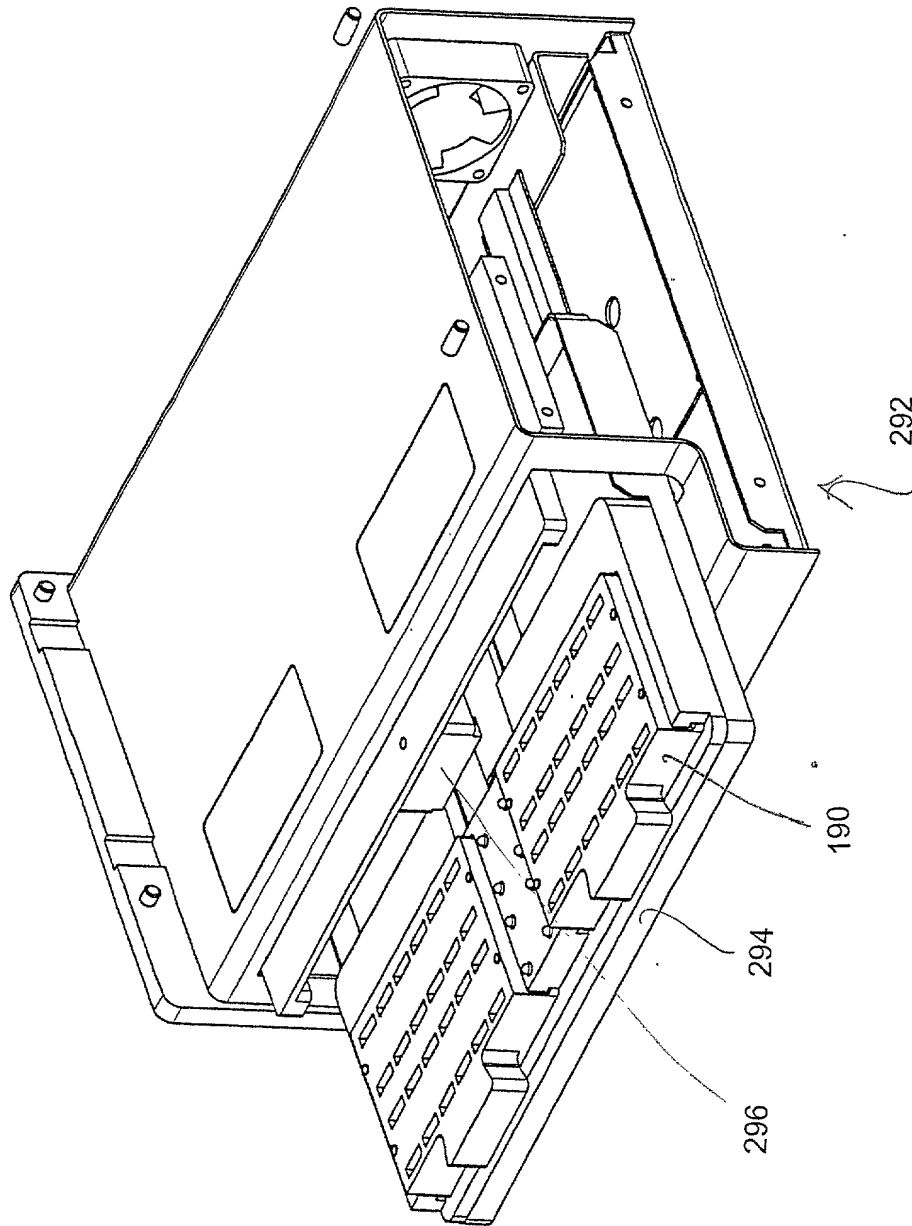


FIG. 13

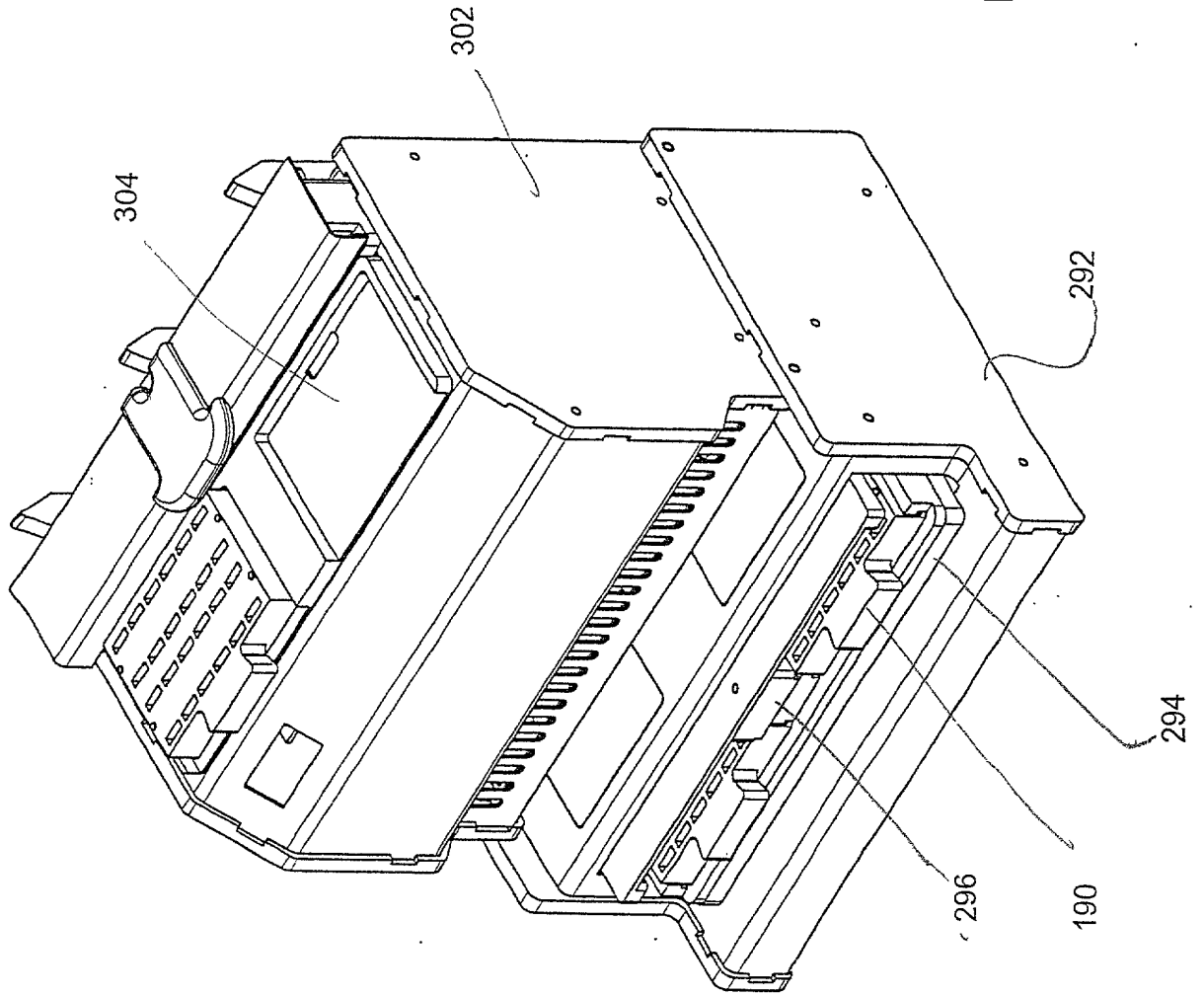


FIG. 14

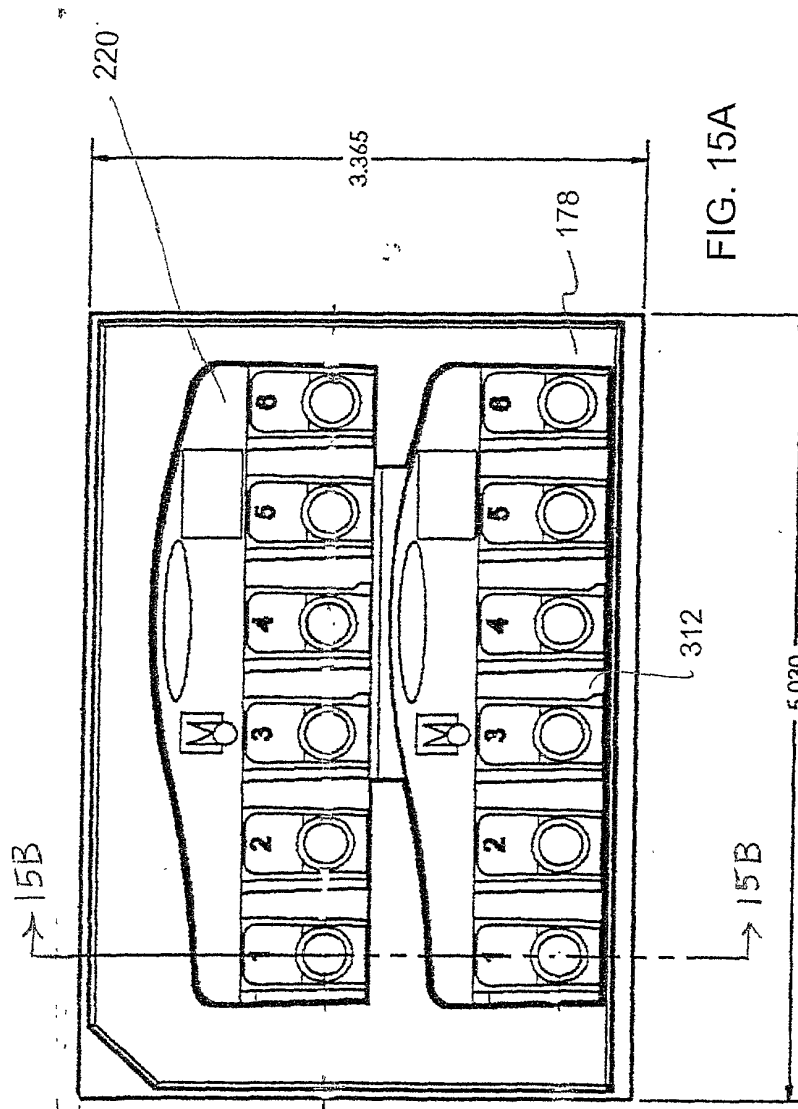


FIG. 15A

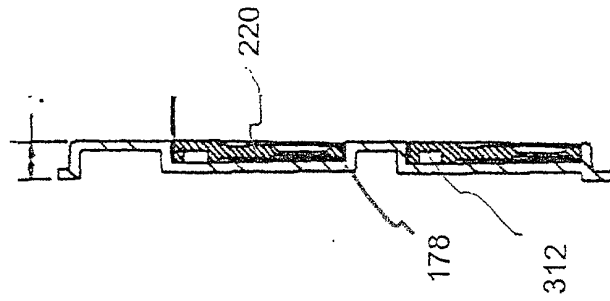


FIG. 15B

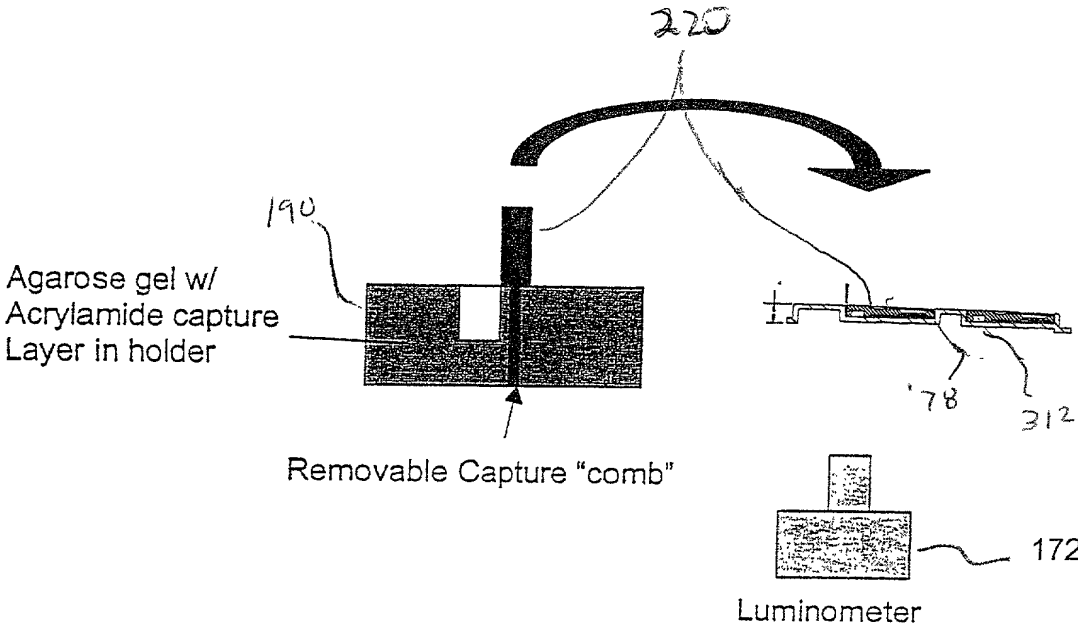


FIG. 16

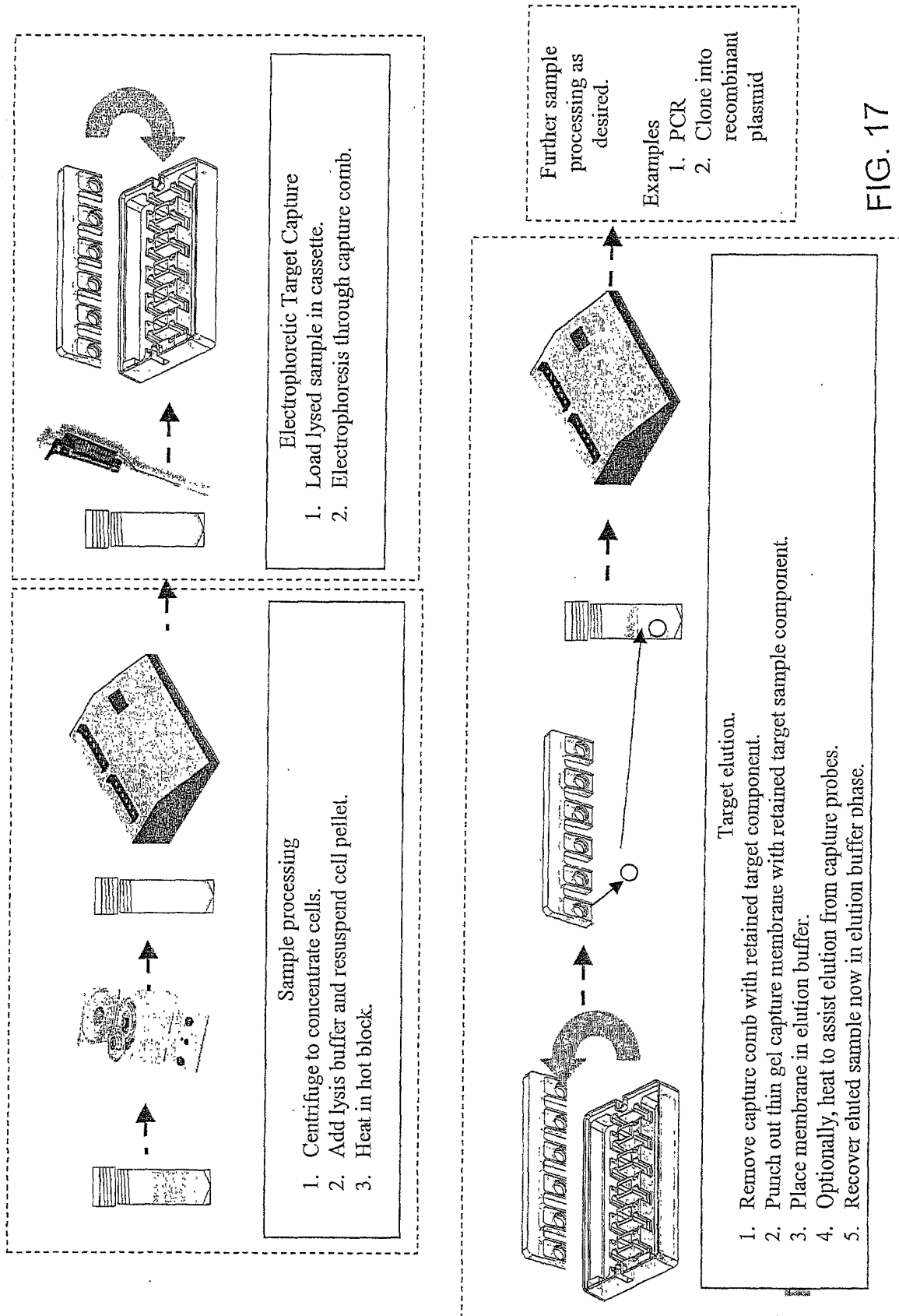


FIG. 17

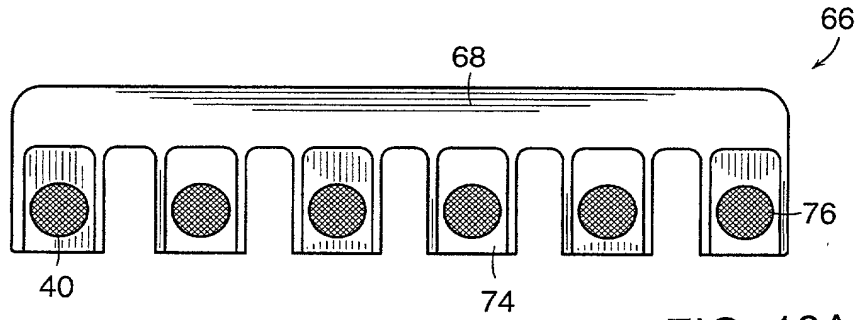


FIG. 18A

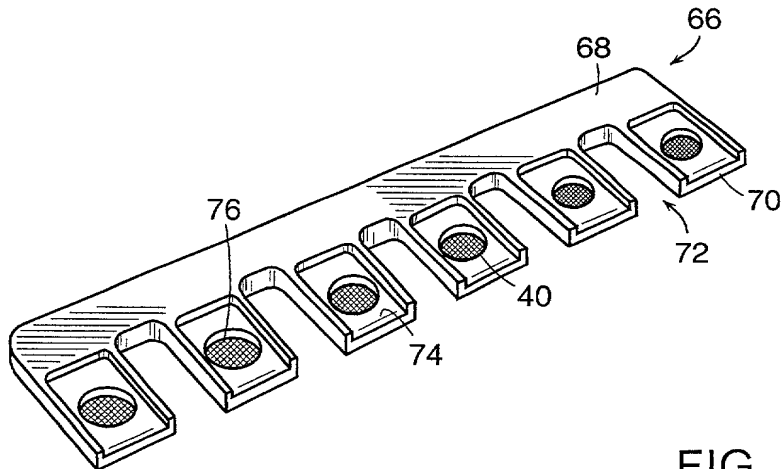


FIG. 18B

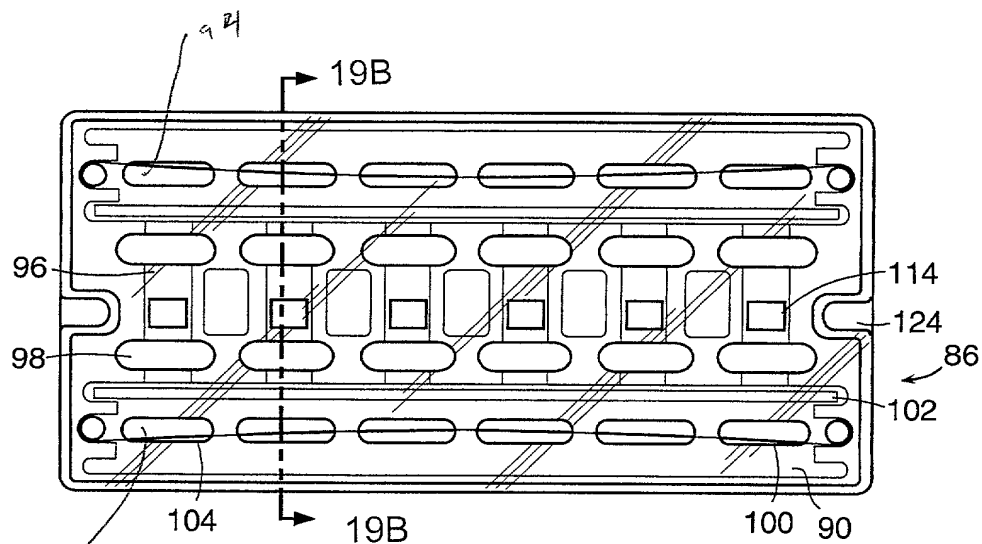


FIG. 19A

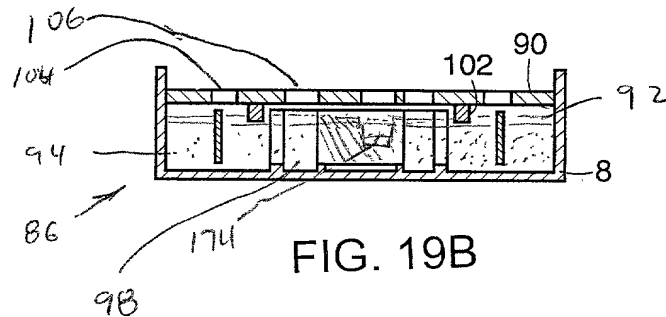


FIG. 19B

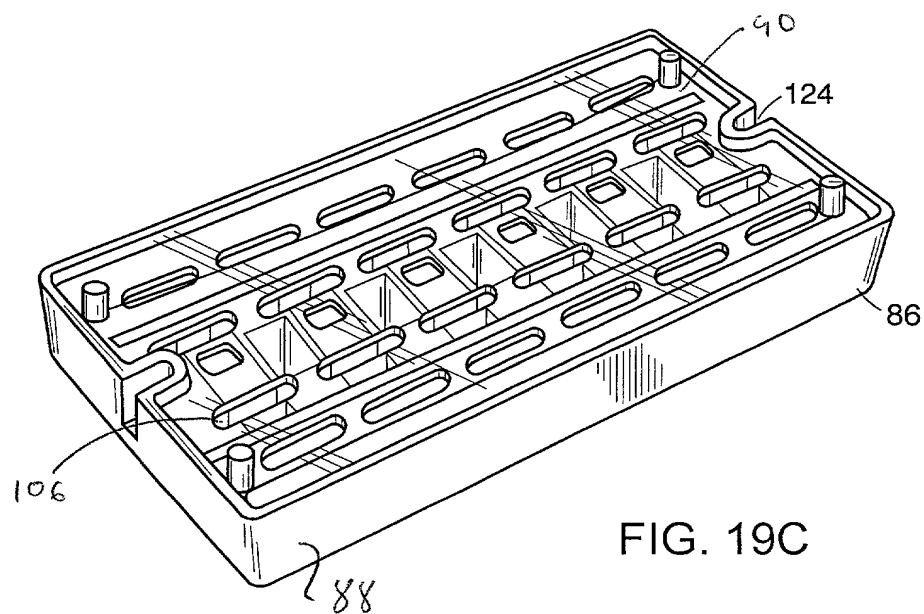


FIG. 19C

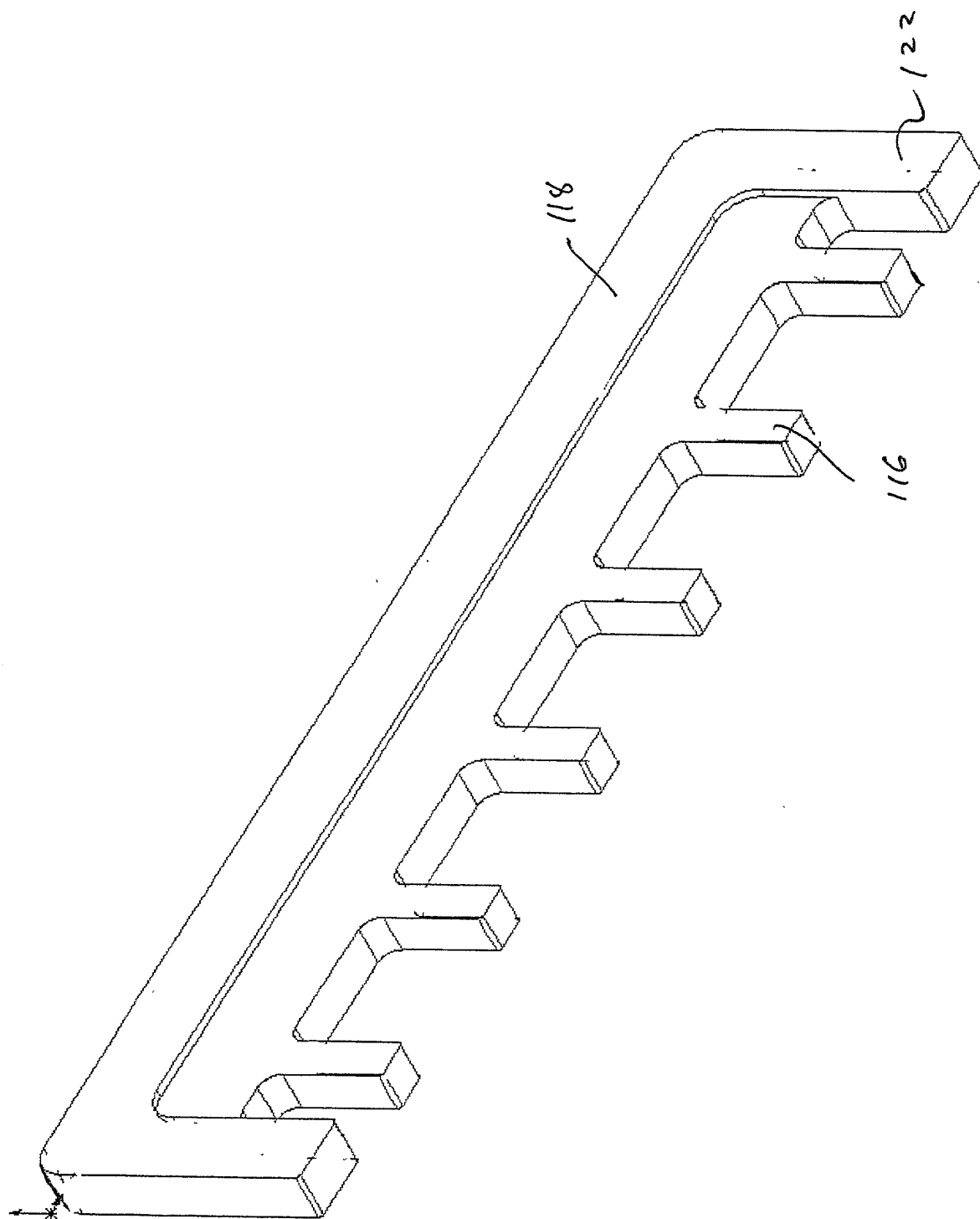
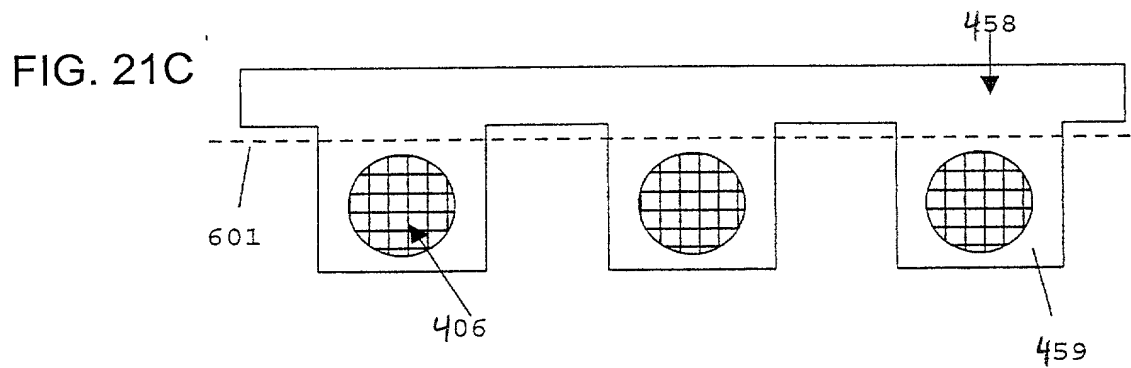
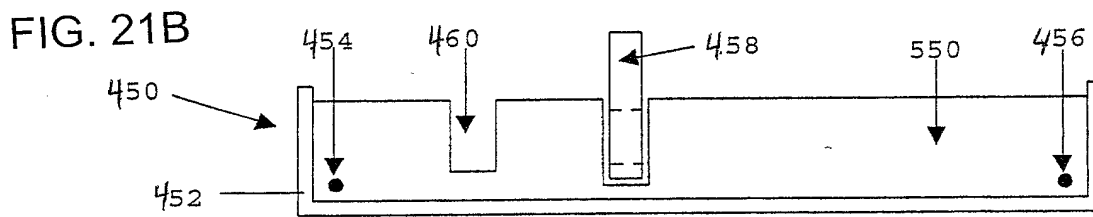
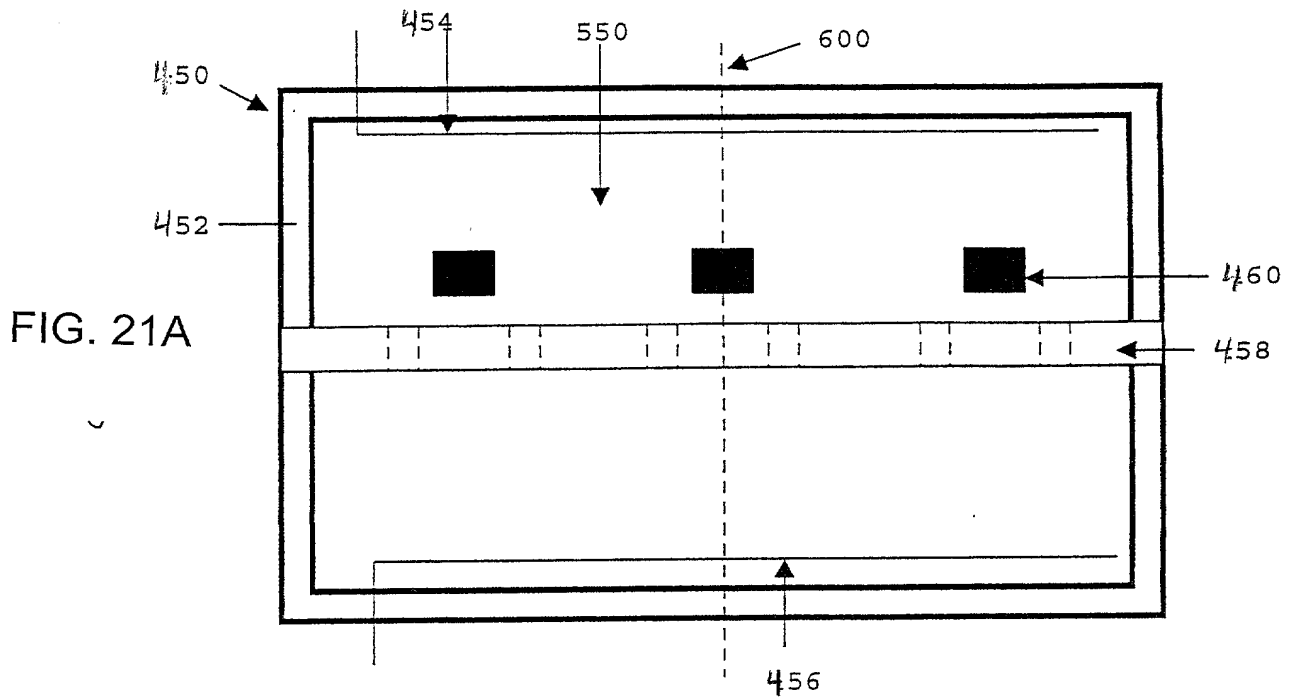


FIG. 20



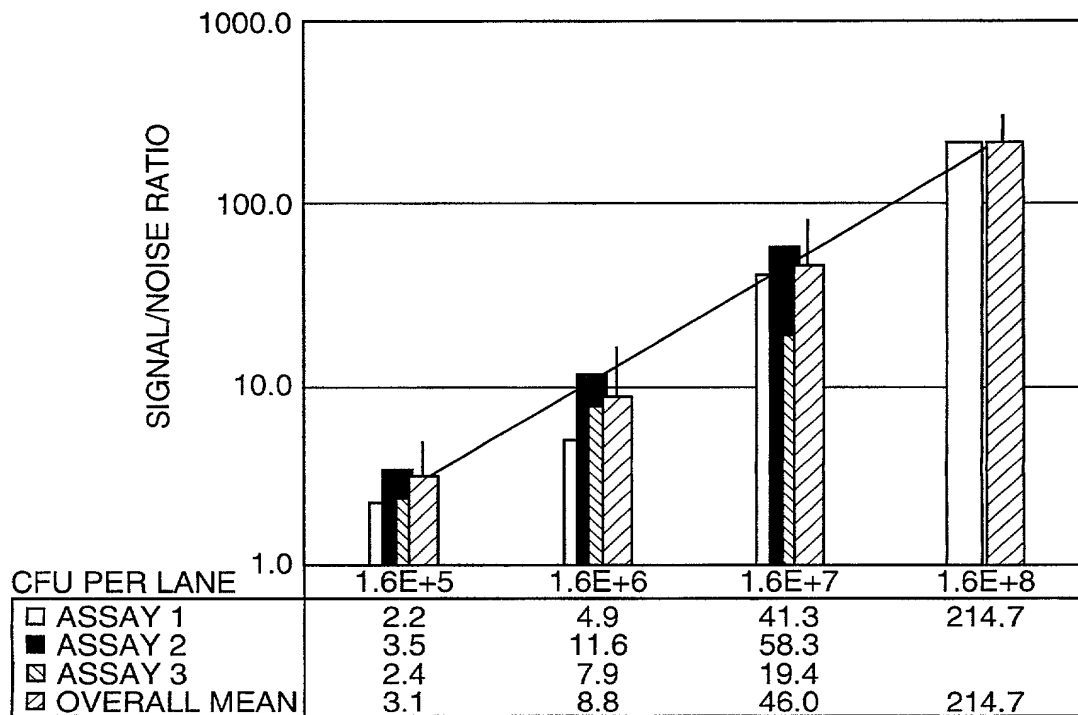
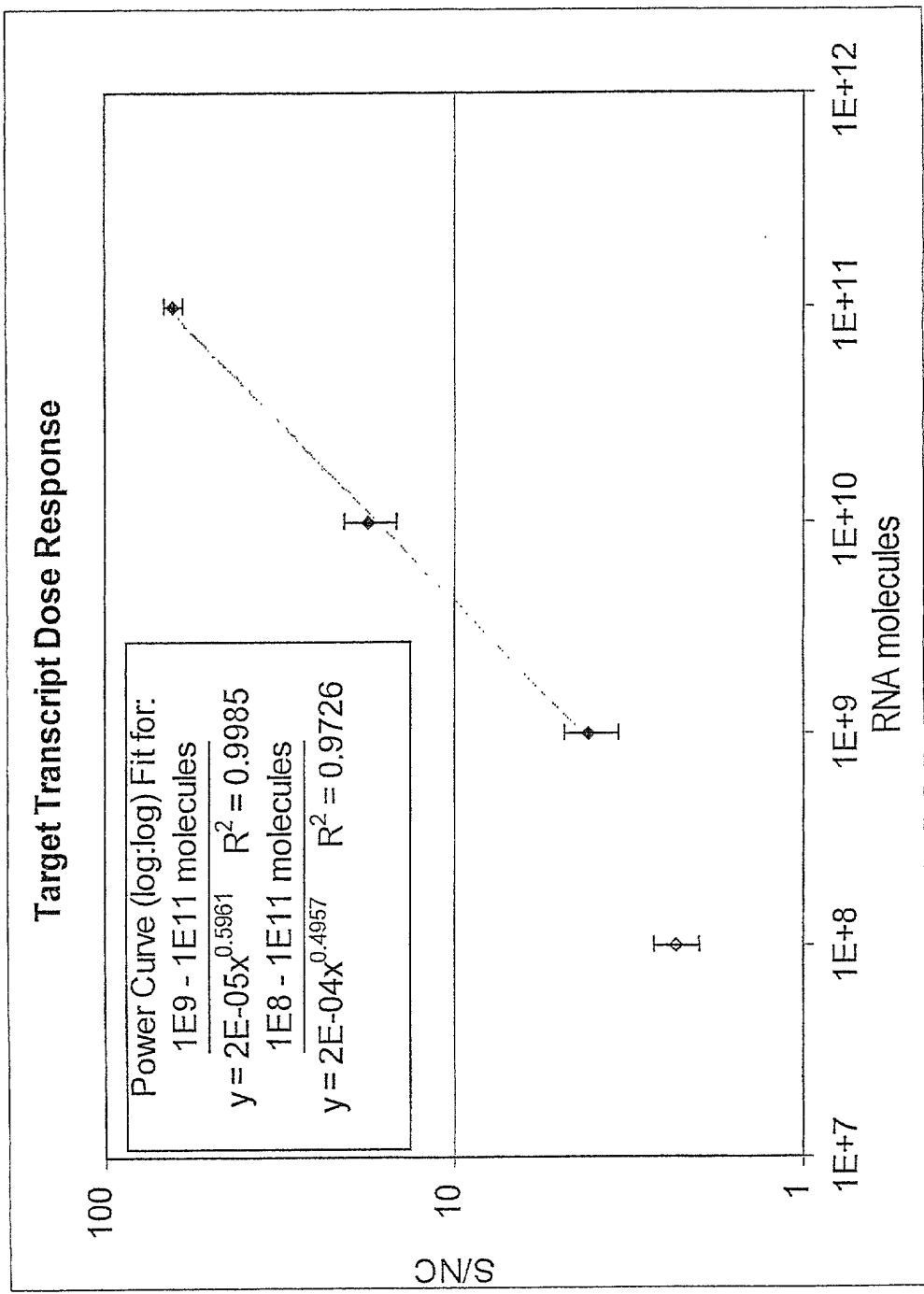
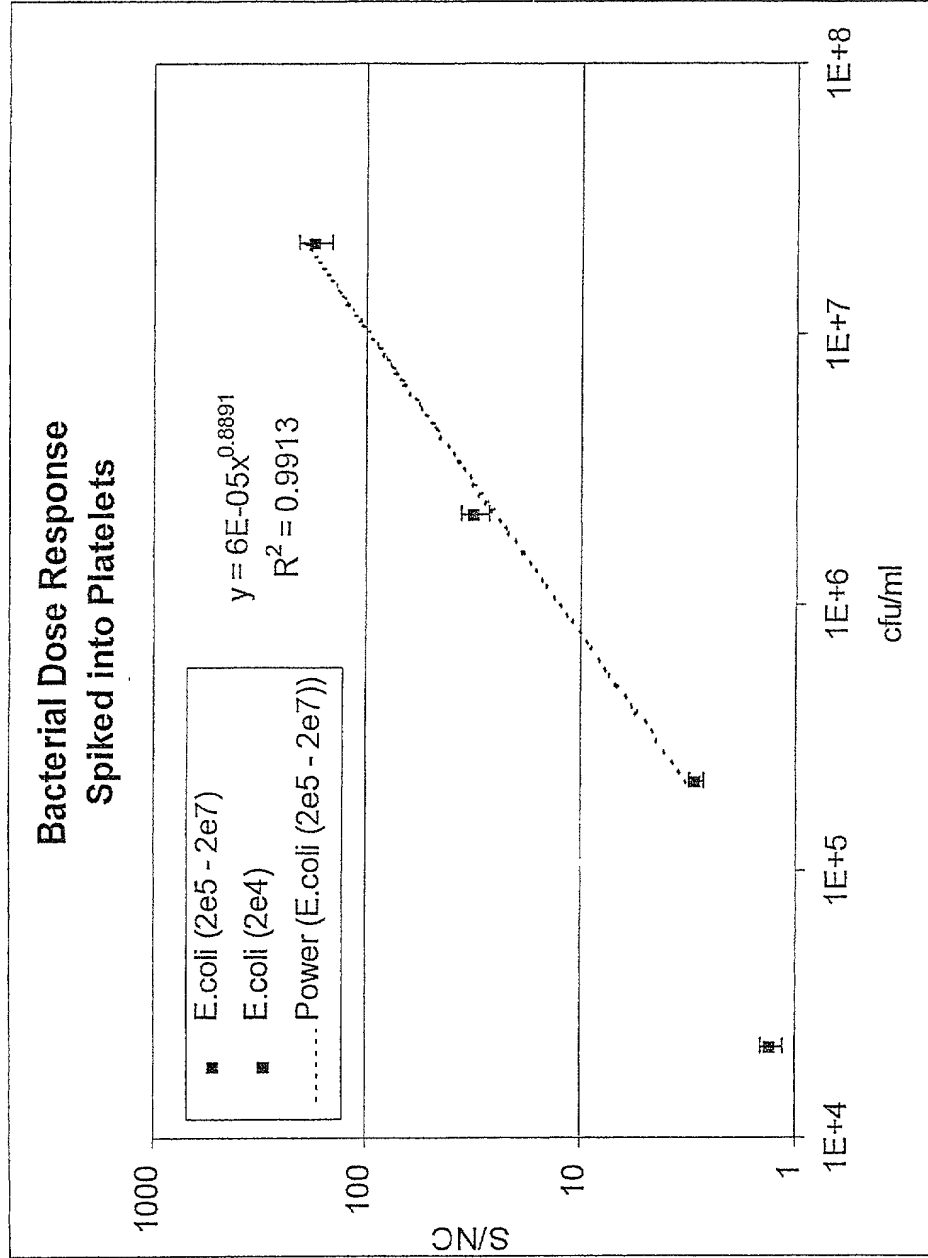


FIG. 22



- S/N from target transcripts spiked into assay after centrifugation step (mean +/- s.e. (n=13))
- Copy# of 4.5S in *E.coli* ~1,000/cfu

FIG. 23



- S/N from *E.coli* (n=8) spiked into negative platelets, then processed according to the assay scheme (mean +/- s.e.).

FIG. 24

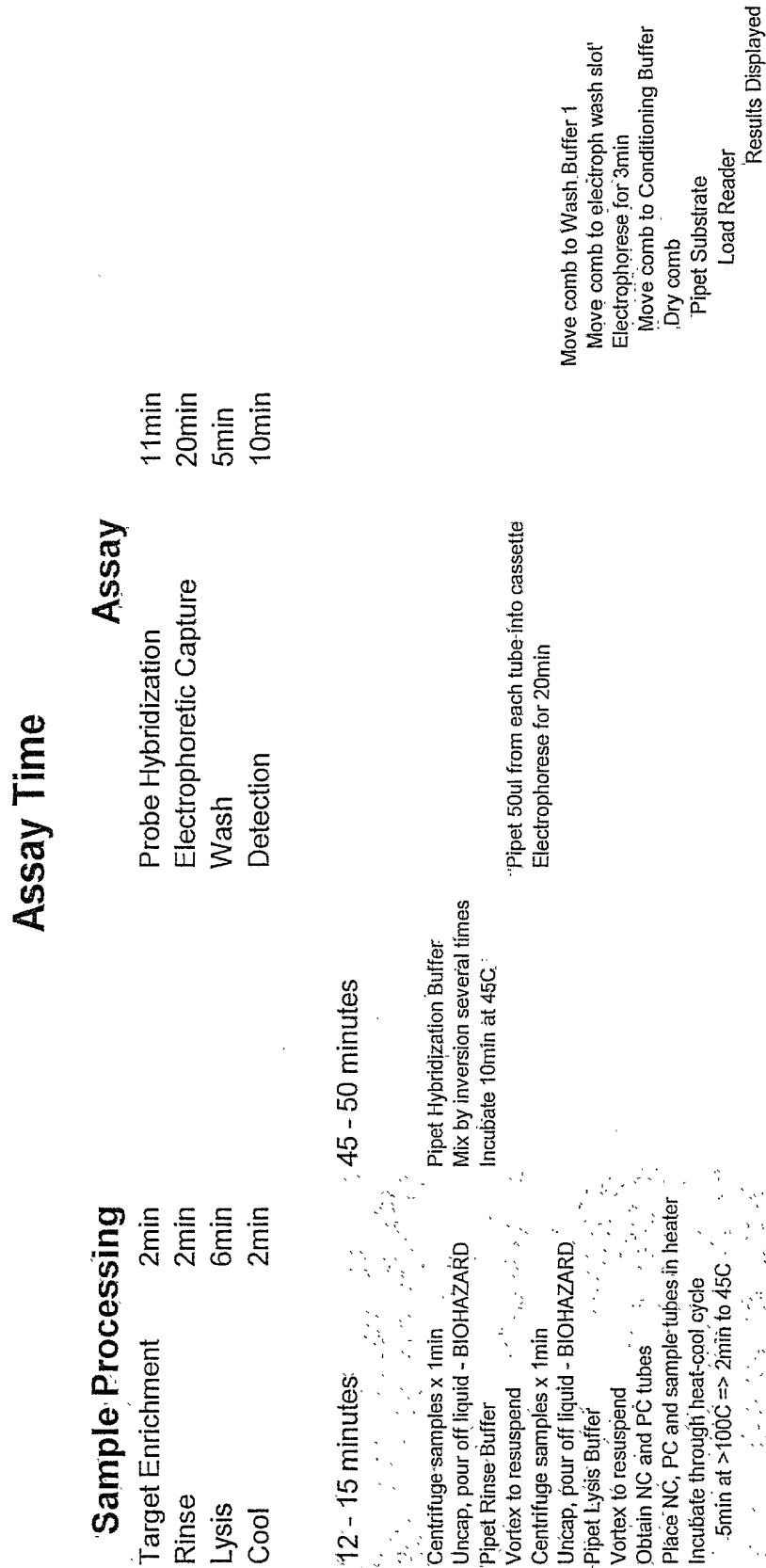


FIG. 25